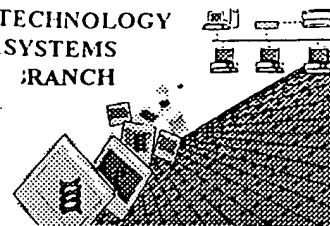


J/PE

0570  
0808BIOTECHNOLOGY  
SYSTEMS  
RANCH

## CRF Problem Report

The Scientific and Technical Information Center (STIC) experienced a problem when processing the following computer readable form (CRF):

Application Serial Number: 09/818583

Filing Date: 03/28/01

Date Processed by STIC: 10/05/01

STIC Contact: Mark Spencer, 703-308-4212

### Nature of Problem:

The CRF (was):

☐ (circle one) Damaged or Unreadable (for Unreadable, see attached)

☐ Blank (no files on CRF) (see attached)

☐ Empty file (filename present, but no bytes in file) (see attached)

☐ Virus-infected. Virus name: \_\_\_\_\_ The STIC will not process the CRF.

☐ Not saved in ASCII text

☐ Sequence Listing was embedded in the file. According to Sequence Rules, submitted file should **only** be the Sequence Listing.

☒ Did not contain a Sequence Listing. (see attached sample)

☐ Other: \_\_\_\_\_

**PLEASE USE THE CHECKER VERSION 3.0 PROGRAM TO REDUCE ERRORS.  
SEE BELOW FOR DETAILS:**

### **Checker Version 3.0**

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 - 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO).

Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

**Checker Version 3.0 can be down loaded from the USPTO website at the following address:**

**<http://www.uspto.gov/web/offices/pac/checker>**

BEST AVAILABLE COPY

09/8/8583

yy`CProject`=There must be an individual or organizational applicant  
name.yy`CProjectData`%Method For Synthesis of Nucleic Acids;P107424-00025  
09/818,583  
2001-03-28`yy`CDNASequence`SEQ ID NO: 1ZArtificial SequenceIgaagagccaa ggacaggtac  
20`flagctrymkswbdhvnE'S`DNA`yy`fl`CCommentFeature`y  
yyyy`1`COtherFeature`misc\_feature,Primer GH20`20`1`SEQ ID NO:  
2ZArtificial SequenceIggaaaataga ccaataggca g  
21`flagctrymkswbdhvn  
e`DNA`yyyy`misc\_feature,Primer  
GH21`21`1`yy`fl`CIndivApplicant`  
`Hiroshi`Tonoike`

Does Not Comply  
Corrected Diskette Needed

Printed File Contents as of  
11:11am 10/05/2001 Marianne Harper

From Diskette  
No sequence looking in diskette  
11:11 am  
10/05/2001  
msh

09/8/8, 583

SEQ ID NO: 2žArtificial SequenceIggaaaataga ccaataggca g  
21~~~~~flagctrymkswbdhvn e ~DNA~~~~~?

SEQ ID NO: 1ŽArtificial SequenceIgaagagccaa ggacaggtag  
20~~~~~flagctrymkswbdhvnE'S~DNA~~~~~yy`n`CCommentFeature`~`  
~~~~~yyyyyy`~`  
`COtherFeature`~`

misc\_feature

```
Primer GH20^20'1'~~~~~?^~~~
```

ÿÿ'^`CPProject'^`=There must be an individual or organizational applicant  
name.ÿÿ'^`

```
`CProjectData`$$$Method For Synthesis of Nucleic Acids$P107424-00025  
09/818,583  
2001-03-28`$$$$$$`yy`
```

`CDNASequence~`~`~`